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Timestamp: [year=2010; month=12; day=28; hr=9; min=30; sec=49; ms=138;]

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Application No: 10728323 Version No: 3.0

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Started: 2010-12-14 14:04:38.691

Finished: 2010-12-14 14:04:39.273

Elapsed: 0 hr(s) 0 min(s) 0 sec(s) 582 ms

Total Warnings: 0

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No. of SeqIDs Defined: 84

Actual SeqID Count: 84

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<110> Bannon, et al.

<120> Microbial Delivery System

<130> 2006517-0010

<140> 10728323

<141> 2003-12-04

<150> 09/731,375

<151> 2000-12-06

<150> 60/195.035

<151> 2000-04-06

<160> 84

<170> PatentIn version 3.5

<210> 1

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<213> Arachis hypogaea

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Thr Thr Asn Gln Arg Ser Pro Pro Gly Glu Arg Thr Arg Gly Arg Gln
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Pro Gly Asp Tyr Asp Asp Asp Arg Arg Gln Pro Arg Arg Glu Glu Gly
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Gly Arg Trp Gly Pro Ala Gly Pro Arg Glu Arg Glu Arg Glu Glu Asp
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Trp Arg Gln Pro Arg Glu Asp Trp Arg Arg Pro Ser His Gln Gln Pro
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Pro Gly Ser His Val Arg Glu Glu Thr Ser Arg Asn Asn Pro Phe Tyr
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195 200 205

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225 230 235 240

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Asn Leu Asp Glu Gly His Ala Leu Arg Ile Pro Ser Gly Phe Ile Ser
260 265 270

Tyr Ile Leu Asn Arg His Asp Asn Gln Asn Leu Arg Val Ala Lys Ile
 275 280 285

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Leu Ala Phe Pro Gly Ser Gly Glu Gln Val Glu Lys Leu Ile Lys Asn
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Gln Lys Glu Ser His Phe Val Ser Ala Arg Pro Gln Ser Gln Ser Gln
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Gln Leu Glu Arg Ala Asn Leu Arg Pro Cys Glu Gln His Leu Met Gln
35 40 45

Lys Ile Gln Arg Asp Glu Asp Ser Tyr Glu Arg Asp Pro Tyr Ser Pro
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Ser Gln Asp Pro Tyr Ser Pro Ser Pro Tyr Asp Arg Arg Gly Ala Gly
65 70 75 80

Ser Ser Gln His Gln Glu Arg Cys Cys Asn Glu Leu Asn Glu Phe Glu
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Asn Asn Gln Arg Cys Met Cys Glu Ala Leu Gln Gln Ile Met Glu Asn
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Gln Ser Asp Arg Leu Gln Gly Arg Gln Gln Glu Gln Gln Phe Lys Arg
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Ala	Leu	Ser	Arg	Leu	Val	Leu	Arg	Arg	Asn	Ala	Leu	Arg	Arg	Pro	Phe	50	55	60	
Tyr	Ser	Asn	Ala	Pro	Gln	Glu	Ile	Phe	Ile	Gln	Gln	Gly	Arg	Gly	Tyr	65	70	75	80
Phe	Gly	Leu	Ile	Phe	Pro	Gly	Cys	Pro	Arg	His	Tyr	Glu	Glu	Pro	His	85	90	95	
Thr	Gln	Gly	Arg	Arg	Ser	Gln	Ser	Gln	Arg	Pro	Pro	Arg	Arg	Leu	Gln	100	105	110	
Gly	Glu	Asp	Gln	Ser	Gln	Gln	Gln	Arg	Asp	Ser	His	Gln	Lys	Val	His	115	120	125	
Arg	Phe	Asp	Glu	Gly	Asp	Leu	Ile	Ala	Val	Pro	Thr	Gly	Val	Ala	Phe	130	135	140	
Trp	Leu	Tyr	Asn	Asp	His	Asp	Thr	Asp	Val	Val	Ala	Val	Ser	Leu	Thr	145	150	155	160
Asp	Thr	Asn	Asn	Asn	Asp	Asn	Gln	Leu	Asp	Gln	Phe	Pro	Arg	Arg	Phe	165	170	175	
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Ser	Arg	Gln	Ser	Arg	Arg	Arg	Ser	Leu	Pro	Tyr	Ser	Pro	Tyr	Ser	Pro	195	200	205	
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Glu Glu Glu Gly Ala Ile Val Thr Val Arg Gly Gly Leu Arg Ile Leu
275 280 285

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Asp Glu Tyr Glu Tyr Asp Glu Glu Asp Arg Arg Arg Gly Arg Gly Ser
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Arg Gly Arg Gly Asn Gly Ile Glu Glu Thr Ile Cys Thr Ala Ser Ala
325 330 335

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Trp Leu Gly Pro Ser Ala Glu Tyr Gly Asn Leu Tyr Arg Asn Ala Leu
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Phe Val Ala His Tyr Asn Thr Asn Ala His Ser Ile Ile Tyr Arg Leu
385 390 395 400

Arg Gly Arg Ala His Val Gln Val Val Asp Ser Asn Gly Asn Arg Val
405 410 415

Tyr Asp Glu Glu Leu Gln Glu Gly His Val Leu Val Val Pro Gln Asn
420 425 430

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Phe Lys Thr Asp Ser Arg Pro Ser Ile Ala Asn Leu Ala Gly Glu Asn

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455

460

Ser Val Ile Asp Asn Leu Pro Glu Glu Val Val Ala Asn Ser Tyr Gly
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